

SEQUENCE PROTOCOL

<110> Degussa-Hüls AG

5 <120> Process for the fermentative preparation of
L-amino acids using strains of the
Enterobacteriaceae family.

<130> 000613 BT

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<170> PatentIn Ver. 2.1

<210> 1

<211> 1719

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<212> DNA

<213> *Escherichia coli*

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<222> (1) .. (1716)

<223> poxB

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ggg gtg aaa cgc atc tgg gga gtc aca ggc gac tct ctg aac ggt ctt 96
Gly Val Lys Arg Ile Trp Gly Val Thr Gly Asp Ser Leu Asn Gly Leu
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40

agt gac agt ctt aat cgc atg ggc acc atc gag tgg atg tcc acc cgc 144
Ser Asp Ser Leu Asn Arg Met Gly Thr Ile Glu Trp Met Ser Thr Arg
35 40 45

cac gaa gaa gtg gcg gcc ttt gcc gct ggc gct gaa gca caa ctt agc 192
His Glu Glu Val Ala Ala Phe Ala Ala Gly Ala Glu Ala Gln Leu Ser
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1003416-1902
205720-244500

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		Glu Leu Val Ser Ser Pro Glu Gln Ile Pro Gln Val Leu Ala Ile Ala	
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5	tcg tca ggt ttc cat acc atg atg aac gcc gac acg tta gtg cta ctc	816
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5 gat ggc ggt ttt agc atg ttg atg ggc gat ttc ctc tca gta gtg cag 1344
Asp Gly Gly Phe Ser Met Leu Met Gly Asp Phe Leu Ser Val Val Gln
435 440 445

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Phe Val Ala Met Glu Met Lys Ala Gly Gly Tyr Leu Thr Asp Gly Thr
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20 gaa cta cac gac aca aac ttt gcc cgc att gcc gaa gcg tgc ggc att 1488
Glu Leu His Asp Thr Asn Phe Ala Arg Ile Ala Glu Ala Cys Gly Ile
485 490 495

25 acg ggt atc cgt gta gaa aaa gcg tct gaa gtt gat gaa gcc ctg caa 1536
Thr Gly Ile Arg Val Glu Lys Ala Ser Glu Val Asp Glu Ala Leu Gln
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30 cgc gcc ttc tcc atc gac ggt ccg gtg ttg gtg gat gtg gtg gtc gcc 1584
Arg Ala Phe Ser Ile Asp Gly Pro Val Leu Val Asp Val Val Val Ala
515 520 525

35 aaa gaa gag tta gcc att cca ccg cag atc aaa ctc gaa cag gcc aaa 1632
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530 535 540

40 ggt ttc agc ctg tat atg ctg cgc gca atc atc agc gga cgc ggt gat 1680
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gaa gtg atc gaa ctg gcg aaa aca aac tgg cta agg taa 1719
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His Glu Glu Val Ala Ala Phe Ala Ala Gly Ala Glu Ala Gln Leu Ser
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15 Gly Glu Leu Ala Val Cys Ala Gly Ser Cys Gly Pro Gly Asn Leu His
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Leu Ile Asn Gly Leu Phe Asp Cys His Arg Asn His Val Pro Val Leu
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Ala Ile Ala Ala His Ile Pro Ser Ser Glu Ile Gly Ser Gly Tyr Phe
100 105 110
25 Gln Glu Thr His Pro Gln Glu Leu Phe Arg Glu Cys Ser His Tyr Cys
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Tyr His Ala Pro Gln Pro Val Val Thr Pro Glu Glu Glu Glu Leu Arg
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10034416.021200

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Ile Gln Ile Asp Ile Asn Pro Ala Ser Ile Gly Ala His Ser Lys Val
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Phe Ala Ala Asp Asp Ala Ile Phe Thr Cys Asp Val Gly Thr Pro Thr
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40 Asp Gly Gly Phe Ser Met Leu Met Gly Asp Phe Leu Ser Val Val Gln
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Met Lys Leu Pro Val Lys Ile Val Val Phe Asn Asn Ser Val Leu Gly
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Arg Ala Phe Ser Ile Asp Gly Pro Val Leu Val Asp Val Val Val Ala
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15 Lys Glu Glu Leu Ala Ile Pro Pro Gln Ile Lys Leu Glu Gln Ala Lys
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30 <213> *Escherichia coli*

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35 <223> Mutagenic DNA

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40 <223> Technical DNA/residues of the polylinker sequence

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 15 <223> 5' region of the delta poxB allele

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 25 <223> 3' region of the delta poxB allele

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 taaggtaa 1448

15

100% of the 1000